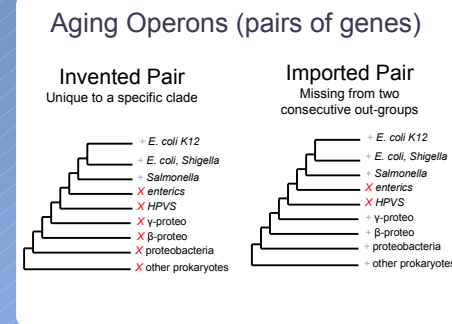


Abstract

Why Are There Operons?

- Co-regulation selects for new operons
 - Operons have more upstream regulatory information
- Operons are not selfish
 - Operons form without horizontal gene transfer (HGT)
 - Non-HGT genes are mostly in operons
 - Non-HGT and native genes form new operons
- New operons are not functionally coherent
- HGT does help maintain operons
 - Associated with transfer of existing operons, not with the creation of new operons
 - Insertion of new genes often creates operons

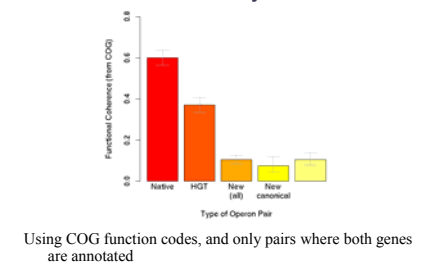


Aging Genes

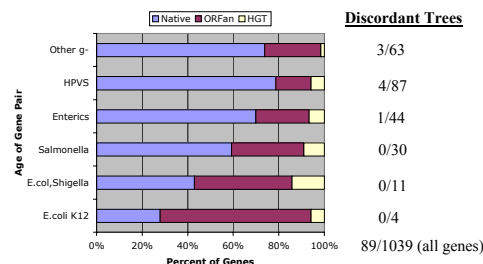
Similar to Aging Operons

- Native to the proteobacteria
 - Homologs in every clade
- Non-HGT (special subset of Native)
 - Single-copy ubiquitous COGs
 - Gene tree agrees with species tree (Lerat et al. 2003)
- ORFan
 - No homologs outside clade

New Operons Are Not as Functionally Coherent



Inventing Operons without HGT

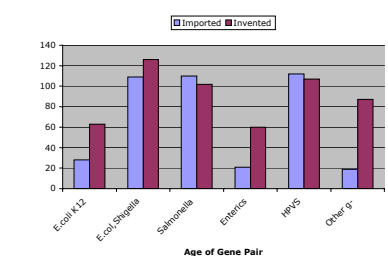


New Operons of Non-HGT Genes

- 37 new operons from 200 non-HGT genes
 - Known operons, or predictions confirmed by microarray data
- Many essential genes (*)
- Sometimes merge two pre-existing operons or add to one

Age of Operon	Known	Microarray	Upstream Gene	Downstream Gene	Age of non-hold gene
Salmonella	No	0.51	yhlB	yhlC	Naive
HPVS	No	0.76	yhlD	yhlE	Naive
HPVS	No	0.89	yhlF	yhlG	Naive
HPVS	No	0.73	yhlH	yhlI	Naive
HPVS	No	0.71	yhlJ	yhlK	Naive
HPVS	No	0.67	yhlL	yhlM	Naive
HPVS	No	0.65	yhlN	yhlO	Naive
HPVS	No	0.61	yhlP	yhlQ	Naive
HPVS	Yes	0.61	yhlR	yhlS	Naive
HPVS	No	0.58	yhlT	yhlU	Naive
HPVS	No	0.54	yhlV	yhlW	Naive
HPVS	Yes	0.13	yhlX	yhlY	Naive
gammata	Yes	0.81	yhlZ	yhlA	Naive
gammata	Yes	0.76	yhlB	yhlC	Naive
gammata	No	0.74	yhlD	yhlE	Naive
gammata	No	0.73	yhlF	yhlG	Naive
gammata	No	0.69	yhlH	yhlI	Naive
gammata	Yes	0.59	yhlJ	yhlK	Naive
gammata	No	0.54	yhlL	yhlM	Naive
gammata	Yes	0.34	yhlN	yhlO	Naive
betagamma	Yes	0.63	yhlP	yhlQ	Naive
betagamma	Yes	N/A	yhlR	yhlS	Naive
betagamma	Yes	0.92	yhlT	yhlU	Naive
betagamma	Yes	0.93	yhlV	yhlW	Naive
betagamma	Yes	0.87	yhlX	yhlY	Naive
betagamma	Yes	0.85	yhlZ	yhlA	Naive
betagamma	Yes	0.72	yhlB	yhlC	Naive
betagamma	No	0.69	yhlD	yhlE	Naive
betagamma	No	0.64	yhlF	yhlG	Naive
betagamma	No	0.54	yhlH	yhlI	Naive
proteo	Yes	0.98	yhlJ	yhlK	Naive
proteo	Yes	0.85	yhlL	yhlM	Naive
proteo	No	0.83	yhlN	yhlO	Naive
proteo	No	0.76	yhlP	yhlQ	Naive
proteo	No	0.74	yhlR	yhlS	Naive
proteo	Yes	0.73	yhlT	yhlU	Naive
proteo	Yes	0.64	yhlV	yhlW	Naive
proteo	Yes	0.63	yhlX	yhlY	Naive
proteo	No	0.52	yhlZ	yhlA	Naive

Many Operons Are Imported



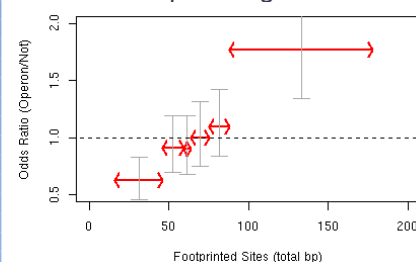
- Imported operons are 2x enriched in HGT genes
- Gene's age usually agrees with operon's age (simple transfer)

Theories of Operon Evolution

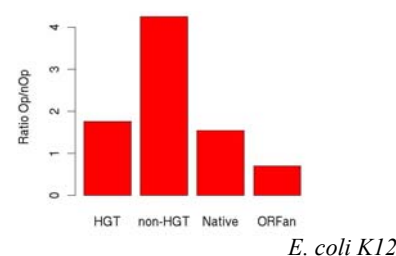
- Co-regulation
 - Clustering genes under the same promoter provides a selective advantage to the organism
- Selfish operons
 - Clustering of genes into operons makes them more likely to propagate by HGT
- Others: Thermoadaptation, Coevolution



Operons Have More Complex Regulation

Phylogenetic footprints in *E. coli*
(from McCue et al. 2002)Only the first gene in predicted operons
Only genes with footprints

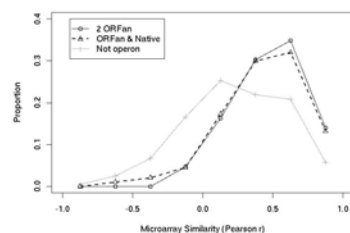
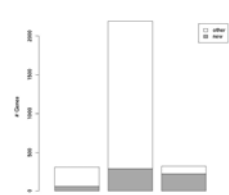
Non-HGT Genes Are More Likely to Be in Operons



Operons Sometimes Form When ORFans Insert

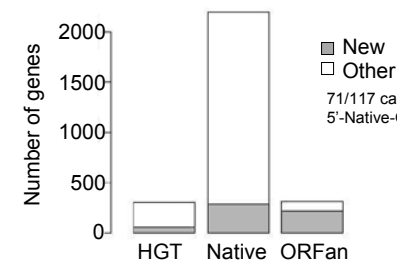
Newest operons* often contain ORFans

Validation of new operons containing ORFans

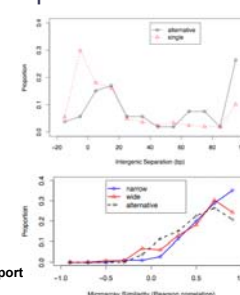


* Unique to Enterobacteria

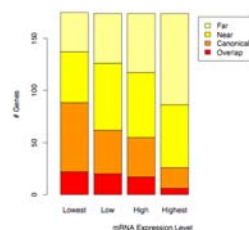
New Operons Contain Many ORFans



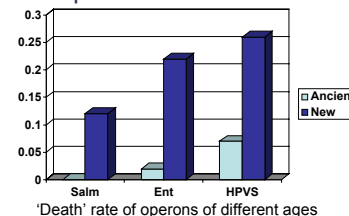
Alternative Transcripts and Widely Spaced Operons Are Similar

47/156 'wide' operons have conserved footprints
Most have support from literature

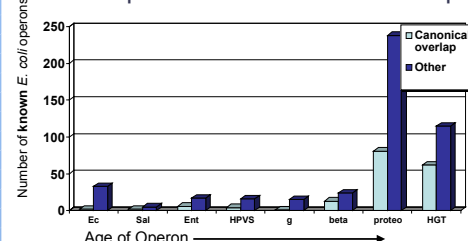
Highly Expressed Genes Have Longer Spacings



New Operons are Less Stable



New Operons Have Fewer Overlaps



Life Cycle of an Operon

- Birth**
 - Gene pairs are merged into new operons
 - By rearrangement or deletion of intervening genes
 - Ancient (and non-HGT) genes can form operons
 - Useful operons are selected for advantageous regulation
 - Some new operons allow ORFans to be expressed?
- Death**
 - Gene order is shuffled extensively across species
 - Operons involving highly conserved or essential genes resist disruption
 - Genes in operons tend to be more conserved & more tightly regulated
 - Immigration
 - Many operons are imported from distant lineages
 - Operons are a useful unit of HGT

